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16. _ 915
/note="platelet membrane glycoprotein; monoclonal antibody /note="platelet membrane glycoprotein; monoclonal antibody F11 induces secretion and aggregation of human platelets by recognizing the F11 receptor; Method: conceptual translation with partial peptide sequencing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /trainiation="MGTKAQVERKLLCLFILAILLCSLALGSVTVHSSEPEVRIPENN PVKLSCAYSGFSSPRVEWKFDQGDTTRLVCYNNKITASYEDRVTFLPTGITFKSVTRE DTGTYTCMVSEEGGNSYGEKKKKLITJVPPSKRTVNIPSSATIGNRATJTCSEQDGSPPSTTWFKDGIVMFTNPKSTRAFSNSSYVLNPTTGELVFDPLSASDTGBYSCEARNGYGTPWTSANKBAVERVYTVAAVLVTLILLGILVFGIWFAYSRGHFDRTKKGTSSKKVIYSQPSARSEGEFKQTSSPLV"
                                                                                                                                                                                                                                                                                                                                                                                                                     Health
NY 11203,
                                                         Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo. (base 100 to 175)
Naik, U. P., Ehrlich, Y. H. and Kornecki, E. Mechanisms of platelet activation-by, a stimulatory antibody: Cross-linking of a novel platelet receptor for monoclonal antibody: Pll with the Fc gamma RII receptor. Biochem. J. 310 (Pt 1), 155-162 (1995)
                                                                                                                                                                                                      2 (bases 1 to 1822)
Sobocka,M.B., Sobocki,T., Banerjee,P., Weiss,C., Rushbrook,J.I.,
Norin,A.J., Hartwig,J., Salifu,M.O., Markell,M.S., Babinska,A.,
Ehrlich,Y.H. and Kornecki,E.
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Sobocka, M.B., Sobocki, T., Rushbrook, J.I., Banerjee, P., Weiss, C.
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                                                                                                                                                                                                                                                                 Cloning of the human platelet F11 receptor: a cell adhesion molecule member of the immunoglobulin superfamily involved in platelet aggregation
Blood 95 (8), 2600-2609 (2000)
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Submitted (19-NOV-1999) Anatomy
Science Center at Brooklyn, 450
USA
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I (bases 1 to 1842)
Wood, W.I., Gurney, A.L., Goddard, A., Pennica, D., Zheng, J. and Yuan, J.
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CCTGAGAATAATCCTGTGAAGTTGTCCTGTGCCTACTCGGGCTTTTTCTTCTCCCCGTGTG GluTrpLysPheAspGlnGlyAspThrThrArgLeuValCysTyrAsnAsnLysIleThr

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THOMO Sapiens (human)
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GluValLysValLysEverileValLeuValProProSerLysProThrValAsnIlePro
                             SerSerAlaThrIleGlyAsnArgAlaValLeuThrCysSerGluGlnAspGlySerPro
                                                                                                                             ProSerGluTyrThrTrpPheLysAspGlyIleValMetProThrAsnProLysSerThr
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                                     This clone is available from RZDD;

This clone is available from RZDD;

Contact RZDP (customer.servicedrzpd.de) for further information.

This CDS clone is a part of a collection of human full ORF clones

Grant GDS has been cloned incl. stopcodon.

The CDS has been cloned incl. stopcodon.

The CDS has been alsorted into pDONR201 via a BP Clonase(TM)

reaction. Additional sequence has been added in front of the start

codon: att. AAAAAA GCA GCG (ATG).

After the stop codon 3' UTR sequence is present in front of the 3'

att site (ACCCAGCTTTCTT).

Compared to the reference sequence NM_144501 (gi21464106) we did

not find any amino acid exchanges.

Clone distribution: http://www.rzpd.de/products/orfclones/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /procedi id="CAG38543.1"
//procedi id="CAG38543.1"
//procedi id="CAG385450"
//procedi id=10="MGTKAQVERKLICLFILAILLCSLALGSVTVHSSEPEVRIPENN VEXASGASSPYEWKEPOGDTTRLVCYNNKITASYEBSVTFLPGITFKGVTRE DTGTYTCMVSEEGGNSYGFSEPWKTRLOLFILVNINKITASYEBSVTFLPGITFKGVTRE DTGTYTCMVSEEGGNSYGFSVKTAIVLIVPPSKPTVNIPSSATIGNRAVLTCSEQDGSP SEYTWRGATOGIVMPTRRKKSTRARSNSSYVLNPTTGELVFDPLSASDTGSYSCEARNGY GTPMTSNAVRAMAVENNVGYIVAAVLYTLILLGILVFGIWFAYSRGHPDRTKKGTSSK KVIYSQPSARSEGBFKQISSFLVN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       241 GCTTCCTATGAGGACCGGGGACCTTCTTGCCAACTGGTATCACCTTCAAGTCGTGACA 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ArgGluAspThrGlyThrTyrThrCysMetValSerGluGluGlyGlyAsnSerTyrGly 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              41 ProGluAsnAsnProValLysLeuSerCysAlaTyrSerGlyPheSerSerProArgVal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GluTrpLysPheAspGlnGlyAspThrThrArgLeuValCysTyrAsnAsnLysIleThr
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                                                                                                                                                                                                                                                                                                                                                                                                                               organism="Homo sapiens"

mol_type="mRNA"

db_xxef="taxon:9606"

clone="REPD0834D0919D"

clone lib="Human Full ORF Clones Gateway(TM) - RZPD"

lab_host="DH108"
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AGIP9/10,
AGIP29/00, AGIP35/00, AGIP37/00, CO7K14/47, CO7K16/18, C12N1/15, PC
  JP 2001509372-A/1
24-JUL-2001
10-JUL-1998 JP 2000502080
110-JUL-1997 US 60/052186
SHALLEY KANT GUPTA, KODANDARAM PILLARISETTI
CIZALE/09,A61K38/00,A61K39/395,A61K45/00,A61P9/00,
                                                                                                                                                                                                                               C12N1/21,C12N5/10,C12P21/02,C12N15/00,A61K37/02,C12N5/00
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1 (bases 1 to 1140)

Gupta, S.K. and Pillarisetti, K.

Human Fil antigen as cell surface receptor participating in platelet coagulation
Patent: JP 2001509372-A 1 24-JUL-2001;

SMITHKLINE BEECHAM COMP.
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